<u>RAW SEQUENCE LISTING</u>

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) no errors detected.

Application Serial Number: _____ 10/561, 132 _____

Source: _____ JFLWP _____

Date Processed by STIC: _____ 01/04/2006 _____

ENTERED

CRF Errors Edited by the STIC Systems Branch

Serial Number: _10/561, 132_

CRF Edit Date: _01/04/2006_
Edited by: _DA_

____ Realigned nucleic acid/amino acid numbers/text in cases where the sequence text "wrapped" to the next line

____ Corrected the SEQ ID NO. Sequence numbers edited were:

_____

____ Inserted or corrected a nucleic number at the end of a nucleic line. SEQ ID NO's edited:

_____

_ . Deleted: ___ invalid beginning/end-of-file text ; ___ page numbers

____ Inserted mandatory headings/numeric identifiers, specifically:

_____

____ Moved responses to same line as heading/numeric identifier, specifically:

_____

Other: _Inserted Mandatory field Identifier_

_____
_____

IFWP

RAW SEQUENCE LISTING   DATE: 01/04/2006
PATENT APPLICATION: US/10/561,132   TIME: 12:51:37

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\01042006\J561132.raw**

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      4         Connolly, Daniel T
      6 <120> TITLE OF INVENTION: Human G Protein-Coupled Receptor and Modulators
Thereof For the
      7         Treatment of Cardiovascular Disorders
      9 <130> FILE REFERENCE: 60.US2.PCT
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/561,132
C--> 11 <141> CURRENT FILING DATE: 2005-12-16
     11 <150> PRIOR APPLICATION NUMBER: 60/480,046
     12 <151> PRIOR FILING DATE: 2003-06-20
     14 <160> NUMBER OF SEQ ID NOS: 12
     16 <170> SOFTWARE: PatentIn version 3.2
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     20 <212> TYPE: DNA
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     30 gaagaataca ctgttaatat tgagatcagt tttgaaaatg catccttcct ggatcctatc     240
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     42 ggctttcaag aagacctcat gaacacttcc tccgccctct ataggtccta caagaccgac     600
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     54 aatgtgtctt ggcgctatga agaacagcag ttggaaatcc agaacagcag cagattctcg     960
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114 gctttcccaa ctctccaagc catccttgct caggatatcc aggaaaataa ctttgcagag  2760
116 agcttagtga tgacaaccac tgtcagccac aatacgacta tgccattcag gatttcaatg  2820
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122 gacaatgtca cctgtatctg tgaccaccta acatcattct ccatcctcat gtcccctgac  3000
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138 gccatctcgg tcatcacgct gggagccacc cagccccggg aagtctatac gaggaagaat  3480
140 gtctgttggc tcaactggga ggacaccaag gccctgctgg ctttcgccat cccagcactg  3540
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154 cctgtgtttt ctatgagttc tccaatatca aggagattta acaatttgtt tggtaaaaca  3960
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168 <221> NAME/KEY: VARIANT
169 <222> LOCATION: 604
170 <223> OTHER INFORMATION: Polymorphic amino acid Met or Thr
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RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/561,132

DATE: 01/04/2006
TIME: 12:51:37

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\01042006\J561132.raw

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177 <222> LOCATION: 856
178 <223> OTHER INFORMATION: Polymorphic amino acid Thr or Met
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187              20                  25                  30
190 Pro Leu Ser Leu His Glu His Glu Pro Ala Gly Glu Glu Ala Leu Arg
191          35                  40                  45
194 Gln Lys Arg Ala Val Ala Thr Lys Ser Pro Thr Ala Glu Glu Tyr Thr
195      50                  55                  60
198 Val Asn Ile Glu Ile Ser Phe Glu Asn Ala Ser Phe Leu Asp Pro Ile
199 65                  70                  75                  80
202 Lys Ala Tyr Leu Asn Ser Leu Ser Phe Pro Ile His Gly Asn Asn Thr
203              85                  90                  95
206 Asp Gln Ile Thr Asp Ile Leu Ser Ile Asn Val Thr Thr Val Cys Arg
207             100                 105                 110
210 Pro Ala Gly Asn Glu Ile Trp Cys Ser Cys Glu Thr Gly Tyr Gly Trp
211         115                 120                 125
214 Pro Arg Glu Arg Cys Leu His Asn Leu Ile Cys Gln Glu Arg Asp Val
215     130                 135                 140
218 Phe Leu Pro Gly His His Cys Ser Cys Leu Lys Glu Leu Pro Pro Asn
219 145                 150                 155                 160
222 Gly Pro Phe Cys Leu Leu Gln Glu Asp Val Thr Leu Asn Met Arg Val
223             165                 170                 175
226 Arg Leu Asn Val Gly Phe Gln Glu Asp Leu Met Asn Thr Ser Ser Ala
227             180                 185                 190
230 Leu Tyr Arg Ser Tyr Lys Thr Asp Leu Glu Thr Ala Phe Arg Lys Gly
231         195                 200                 205
234 Tyr Gly Ile Leu Pro Gly Phe Lys Gly Val Thr Val Thr Gly Phe Lys
235     210                 215                 220
238 Ser Gly Ser Val Val Val Thr Tyr Glu Val Lys Thr Thr Pro Pro Ser
239 225                 230                 235                 240
242 Leu Glu Leu Ile His Lys Ala Asn Glu Gln Val Val Gln Ser Leu Asn
243             245                 250                 255
246 Gln Thr Tyr Lys Met Asp Tyr Asn Ser Phe Gln Ala Val Thr Ile Asn
247             260                 265                 270
250 Glu Ser Asn Phe Phe Val Thr Pro Glu Ile Ile Phe Glu Gly Asp Thr
251         275                 280                 285
254 Val Ser Leu Val Cys Glu Lys Glu Val Leu Ser Ser Asn Val Ser Trp
255     290                 295                 300
258 Arg Tyr Glu Glu Gln Gln Leu Glu Ile Gln Asn Ser Ser Arg Phe Ser
259 305                 310                 315                 320
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263             325                 330                 335
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271         355             360             365
274 Pro Ile Gln Ile Leu Ala Asn Glu Glu Met Lys Val Met Cys Asp Asn
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279 385             390             395             400
282 Lys Val Glu Trp Lys Gln Glu Gly Lys Ile Asn Ile Pro Gly Thr Pro
283             405             410             415
286 Glu Thr Asp Ile Asp Ser Ser Cys Ser Arg Tyr Thr Leu Lys Ala Asp
287             420             425             430
290 Gly Thr Gln Cys Pro Ser Gly Ser Ser Gly Thr Thr Val Ile Tyr Thr
291         435             440             445
294 Cys Glu Phe Ile Ser Ala Tyr Gly Ala Arg Gly Ser Ala Asn Ile Lys
295     450             455             460
298 Val Thr Phe Ile Ser Val Ala Asn Leu Thr Ile Thr Pro Asp Pro Ile
299 465             470             475             480
302 Ser Val Ser Glu Gly Gln Asn Phe Ser Ile Lys Cys Ile Ser Asp Val
303             485             490             495
306 Ser Asn Tyr Asp Glu Val Tyr Trp Asn Thr Ser Ala Gly Ile Lys Ile
307             500             505             510
310 Tyr Gln Arg Phe Tyr Thr Thr Arg Arg Tyr Leu Asp Gly Ala Glu Ser
311             515             520             525
314 Val Leu Thr Val Lys Thr Ser Thr Arg Glu Trp Asn Gly Thr Tyr His
315         530             535             540
318 Cys Ile Phe Arg Tyr Lys Asn Ser Tyr Ser Ile Ala Thr Lys Asp Val
319 545             550             555             560
322 Ile Val His Pro Leu Pro Leu Lys Leu Asn Ile Met Val Asp Pro Leu
323             565             570             575
326 Glu Ala Thr Val Ser Cys Ser Gly Ser His His Ile Lys Cys Cys Ile
327             580             585             590
330 Glu Glu Asp Gly Asp Tyr Lys Val Thr Phe His Met Gly Ser Ser Ser
331             595             600             605
334 Leu Pro Ala Ala Lys Glu Val Asn Lys Lys Gln Val Cys Tyr Lys His
335     610             615             620
338 Asn Phe Asn Ala Ser Ser Val Ser Trp Cys Ser Lys Thr Val Asp Val
339 625             630             635             640
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343             645             650             655
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347             660             665             670
350 Val Ile Gly Val Gly Glu Pro Gly Lys Val Ile Gln Lys Leu Cys Arg
351         675             680             685
354 Phe Ser Asn Val Pro Ser Ser Pro Glu Ser Pro Ile Gly Gly Thr Ile
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359 705             710             715             720
362 Ile Ser Ala Pro Ile Asn Ser Leu Leu Gln Met Ala Lys Ala Leu Ile
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RAW SEQUENCE LISTING   DATE: 01/04/2006
PATENT APPLICATION: US/10/561,132   TIME: 12:51:37

Input Set : A:\pto.da.txt
Output Set: N:\CRF4\01042006\J561132.raw

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367             740               745               750
370 Ser Ile Ser Ile Asp Lys Ala Glu His Glu Ile Ser Ser Ser Pro Gly
371             755               760               765
374 Ser Leu Gly Ala Ile Ile Asn Ile Leu Asp Leu Leu Ser Thr Val Pro
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378 Thr Gln Val Asn Ser Glu Met Met Thr His Val Leu Ser Thr Val Asn
379 785               790               795               800
382 Val Ile Leu Gly Lys Pro Val Leu Asn Thr Trp Lys Val Leu Gln Gln
383             805               810               815
386 Gln Trp Thr Asn Gln Ser Ser Gln Leu Leu His Ser Val Glu Arg Phe
387             820               825               830
390 Ser Gln Ala Leu Gln Ser Gly Asp Ser Pro Pro Leu Ser Phe Ser Gln
391         835               840               845
394 Thr Asn Val Gln Met Ser Ser Thr Val Ile Lys Ser Ser His Pro Glu
395     850               855               860
398 Thr Tyr Gln Gln Arg Phe Val Phe Pro Tyr Phe Asp Leu Trp Gly Asn
399 865               870               875               880
402 Val Val Ile Asp Lys Ser Tyr Leu Glu Asn Leu Gln Ser Asp Ser Ser
403             885               890               895
406 Ile Val Thr Met Ala Phe Pro Thr Leu Gln Ala Ile Leu Ala Gln Asp
407         900               905               910
410 Ile Gln Glu Asn Asn Phe Ala Glu Ser Leu Val Met Thr Thr Thr Val
411     915               920               925
414 Ser His Asn Thr Thr Met Pro Phe Arg Ile Ser Met Thr Phe Lys Asn
415     930               935               940
418 Asn Ser Pro Ser Gly Gly Glu Thr Lys Cys Val Phe Trp Asn Phe Arg
419 945               950               955               960
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423             965               970               975
426 Glu Gly Asp Gly Asp Asn Val Thr Cys Ile Cys Asp His Leu Thr Ser
427         980               985               990
430 Phe Ser Ile Leu Met Ser Pro Asp  Ser Pro Asp Pro Ser  Ser Leu Leu
431         995               1000              1005
434 Gly Ile  Leu Leu Asp Ile Ile  Ser Tyr Val Gly Val  Gly Phe Ser
435     1010              1015              1020
438 Ile Leu  Ser Leu Ala Ala Cys  Leu Val Val Glu Ala  Val Val Trp
439     1025              1030              1035
442 Lys Ser  Val Thr Lys Asn Arg  Thr Ser Tyr Met Arg  His Thr Cys
443     1040              1045              1050
446 Ile Val  Asn Ile Ala Ala Ser  Leu Leu Val Ala Asn  Thr Trp Phe
447     1055              1060              1065
450 Ile Val  Val Ala Ala Ile Gln  Asp Asn Arg Tyr Ile  Leu Cys Lys
451     1070              1075              1080
454 Thr Ala  Cys Val Ala Ala Thr  Phe Phe Ile His Phe  Phe Tyr Leu
455     1085              1090              1095
458 Ser Val  Phe Phe Trp Met Leu  Thr Leu Gly Leu Met  Leu Phe Tyr
459     1100              1105              1110
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RAW SEQUENCE LISTING ERROR SUMMARY DATE: 01/04/2006
PATENT APPLICATION: US/10/561,132   TIME: 12:51:38

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\01042006\J561132.raw**

Invalid <213> Response:
Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.

Seq#:7,8,9,10,11,12

VERIFICATION SUMMARY   DATE: 01/04/2006
PATENT APPLICATION: **US/10/561,132**   TIME: 12:51:38

Input Set : **A:\pto.da.txt**
Output Set: **N:\CRF4\01042006\J561132.raw**

L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:176 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:180 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2

Raw Sequence Listing before editing,
for reference only

)FWP

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            RAW SEQUENCE LISTING              DATE: 12/30/2005
            PATENT APPLICATION:  US/10/561,132    TIME: 09:22:52

            Input Set : A:\60.US2.PCT.txt
            Output Set: N:\CRF4\12302005\J561132.raw

    3 <110> APPLICANT: Adams, John W
    4        Connolly, Daniel T
    6 <120> TITLE OF INVENTION: Human G Protein-Coupled Receptor and Modulators Thereof For
the
    7        Treatment of Cardiovascular Disorders
    9 <130> FILE REFERENCE: 60.US2.PCT
C--> 11 <140> CURRENT APPLICATION NUMBER: US/10/561,132
C--> 11 <141> CURRENT FILING DATE: 2005-12-16
    11 <150> PRIOR APPLICATION NUMBER: 60/480,046
    12 <151> PRIOR FILING DATE: 2003-06-20
    14 <160> NUMBER OF SEQ ID NOS: 12
    16 <170> SOFTWARE: PatentIn version 3.2
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Does Not Comply
Corrected Diskette Needed

Cp٩⁻')

ERRORED SEQUENCES

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    162 <211> LENGTH: 1346
    163 <212> TYPE: PRT
    164 <213> ORGANISM: Homo sapien    --> Inserted ⊂220⁷
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    167 <222> LOCATION: 604
    168 <223> OTHER INFORMATION: Polymorphic amino acid Met or Thr
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    172 <223> OTHER INFORMATION: Polymorphic amino acid Val or Ile
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    185             20                  25                  30
    188 Pro Leu Ser Leu His Glu His Glu Pro Ala Gly Glu Glu Ala Leu Arg
    189             35                  40                  45
    192 Gln Lys Arg Ala Val Ala Thr Lys Ser Pro Thr Ala Glu Glu Tyr Thr
    193         50                  55                  60
    196 Val Asn Ile Glu Ile Ser Phe Glu Asn Ala Ser Phe Leu Asp Pro Ile
    197 65                  70                  75                  80
    200 Lys Ala Tyr Leu Asn Ser Leu Ser Phe Pro Ile His Gly Asn Asn Thr
    201             85                  90                  95
    204 Asp Gln Ile Thr Asp Ile Leu Ser Ile Asn Val Thr Thr Val Cys Arg
    205             100                 105                 110
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209         115             120             125
212 Pro Arg Glu Arg Cys Leu His Asn Leu Ile Cys Gln Glu Arg Asp Val
213     130             135             140
216 Phe Leu Pro Gly His His Cys Ser Cys Leu Lys Glu Leu Pro Pro Asn
217 145             150             155             160
220 Gly Pro Phe Cys Leu Leu Gln Glu Asp Val Thr Leu Asn Met Arg Val
221             165             170             175
224 Arg Leu Asn Val Gly Phe Gln Glu Asp Leu Met Asn Thr Ser Ser Ala
225         180             185             190
228 Leu Tyr Arg Ser Tyr Lys Thr Asp Leu Glu Thr Ala Phe Arg Lys Gly
229     195             200             205
232 Tyr Gly Ile Leu Pro Gly Phe Lys Gly Val Thr Val Thr Gly Phe Lys
233     210             215             220
236 Ser Gly Ser Val Val Val Thr Tyr Glu Val Lys Thr Thr Pro Pro Ser
237 225             230             235             240
240 Leu Glu Leu Ile His Lys Ala Asn Glu Gln Val Val Gln Ser Leu Asn
241             245             250             255
244 Gln Thr Tyr Lys Met Asp Tyr Asn Ser Phe Gln Ala Val Thr Ile Asn
245         260             265             270
248 Glu Ser Asn Phe Phe Val Thr Pro Glu Ile Ile Phe Glu Gly Asp Thr
249         275             280             285
252 Val Ser Leu Val Cys Glu Lys Glu Val Leu Ser Ser Asn Val Ser Trp
253     290             295             300
256 Arg Tyr Glu Glu Gln Gln Leu Glu Ile Gln Asn Ser Ser Arg Phe Ser
257 305         ⁄       310             315                 320
260 Ile Tyr Thr Ala Leu Phe Asn Asn Met Thr Ser Val Ser Lys Leu Thr
261             325             330             335
264 Ile His Asn Ile Thr Pro Gly Asp Ala Gly Glu Tyr Val Cys Lys Leu
265         340             345             350
268 Ile Leu Asp Ile Phe Glu Tyr Glu Cys Lys Lys Lys Ile Asp Val Met
269         355             360             365
272 Pro Ile Gln Ile Leu Ala Asn Glu Glu Met Lys Val Met Cys Asp Asn
273     370             375             380
276 Asn Pro Val Ser Leu Asn Cys Cys Ser Gln Gly Asn Val Asn Trp Ser
277 385             390             395                 400
280 Lys Val Glu Trp Lys Gln Glu Gly Lys Ile Asn Ile Pro Gly Thr Pro
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284 Glu Thr Asp Ile Asp Ser Ser Cys Ser Arg Tyr Thr Leu Lys Ala Asp
285         420             425             430
288 Gly Thr Gln Cys Pro Ser Gly Ser Ser Gly Thr Thr Val Ile Tyr Thr
289         435             440             445
292 Cys Glu Phe Ile Ser Ala Tyr Gly Ala Arg Gly Ser Ala Asn Ile Lys
293     450             455             460
296 Val Thr Phe Ile Ser Val Ala Asn Leu Thr Ile Thr Pro Asp Pro Ile
297 465             470             475                 480
300 Ser Val Ser Glu Gly Gln Asn Phe Ser Ile Lys Cys Ile Ser Asp Val
301             485             490             495
304 Ser Asn Tyr Asp Glu Val Tyr Trp Asn Thr Ser Ala Gly Ile Lys Ile
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308 Tyr Gln Arg Phe Tyr Thr Thr Arg Arg Tyr Leu Asp Gly Ala Glu Ser
309        515                520                525
312 Val Leu Thr Val Lys Thr Ser Thr Arg Glu Trp Asn Gly Thr Tyr His
313        530                535                540
316 Cys Ile Phe Arg Tyr Lys Asn Ser Tyr Ser Ile Ala Thr Lys Asp Val
317 545                550                555                560
320 Ile Val His Pro Leu Pro Leu Lys Leu Asn Ile Met Val Asp Pro Leu
321                565                570                575
324 Glu Ala Thr Val Ser Cys Ser Gly Ser His His Ile Lys Cys Cys Ile
325        580                585                590
328 Glu Glu Asp Gly Asp Tyr Lys Val Thr Phe His Met Gly Ser Ser Ser
329        595                600                605
332 Leu Pro Ala Ala Lys Glu Val Asn Lys Lys Gln Val Cys Tyr Lys His
333        610                615                620
336 Asn Phe Asn Ala Ser Ser Val Ser Trp Cys Ser Lys Thr Val Asp Val
337 625                630                635                640
340 Cys Cys His Phe Thr Asn Ala Ala Asn Asn Ser Val Trp Ser Pro Ser
341                645                650                655
344 Met Lys Leu Asn Leu Val Pro Gly Glu Asn Ile Thr Cys Gln Asp Pro
345        660                665                670
348 Val Ile Gly Val Gly Glu Pro Gly Lys Val Ile Gln Lys Leu Cys Arg
349        675                680                685
352 Phe Ser Asn Val Pro Ser Ser Pro Glu Ser Pro Ile Gly Gly Thr Ile
353        690                695                700
356 Thr Tyr Lys Cys Val Gly Ser Gln Trp Glu Glu Lys Arg Asn Asp Cys
357 705                710                715                720
360 Ile Ser Ala Pro Ile Asn Ser Leu Leu Gln Met Ala Lys Ala Leu Ile
361                725                730                735
364 Lys Ser Pro Ser Gln Asp Glu Met Leu Pro Thr Tyr Leu Lys Asp Leu
365        740                745                750
368 Ser Ile Ser Ile Asp Lys Ala Glu His Glu Ile Ser Ser Ser Pro Gly
369        755                760                765
372 Ser Leu Gly Ala Ile Ile Asn Ile Leu Asp Leu Leu Ser Thr Val Pro
373        770                775                780
376 Thr Gln Val Asn Ser Glu Met Met Thr His Val Leu Ser Thr Val Asn
377 785                790                795                800
380 Val Ile Leu Gly Lys Pro Val Leu Asn Thr Trp Lys Val Leu Gln Gln
381                805                810                815
384 Gln Trp Thr Asn Gln Ser Ser Gln Leu Leu His Ser Val Glu Arg Phe
385        820                825                830
388 Ser Gln Ala Leu Gln Ser Gly Asp Ser Pro Pro Leu Ser Phe Ser Gln
389        835                840                845
392 Thr Asn Val Gln Met Ser Ser Thr Val Ile Lys Ser Ser His Pro Glu
393        850                855                860
396 Thr Tyr Gln Gln Arg Phe Val Phe Pro Tyr Phe Asp Leu Trp Gly Asn
397 865                870                875                880
400 Val Val Ile Asp Lys Ser Tyr Leu Glu Asn Leu Gln Ser Asp Ser Ser
401                885                890                895
```

RAW SEQUENCE LISTING   DATE: 12/30/2005
PATENT APPLICATION: US/10/561,132   TIME: 09:22:52

Input Set : A:\60.US2.PCT.txt
Output Set: N:\CRF4\12302005\J561132.raw

```
404 Ile Val Thr Met Ala Phe Pro Thr Leu Gln Ala Ile Leu Ala Gln Asp
405             900             905             910
408 Ile Gln Glu Asn Asn Phe Ala Glu Ser Leu Val Met Thr Thr Thr Val
409         915             920             925
412 Ser His Asn Thr Thr Met Pro Phe Arg Ile Ser Met Thr Phe Lys Asn
413         930             935             940
416 Asn Ser Pro Ser Gly Gly Glu Thr Lys Cys Val Phe Trp Asn Phe Arg
417 945                 950             955                 960
420 Leu Ala Asn Asn Thr Gly Gly Trp Asp Ser Ser Gly Cys Tyr Val Glu
421             965             970             975
424 Glu Gly Asp Gly Asp Asn Val Thr Cys Ile Cys Asp His Leu Thr Ser
425             980             985             990
428 Phe Ser Ile Leu Met Ser Pro Asp  Ser Pro Asp Pro Ser  Ser Leu Leu
429         995             1000            1005
432 Gly Ile  Leu Leu Asp Ile Ile  Ser Tyr Val Gly Val  Gly Phe Ser
433     1010            1015            1020
436 Ile Leu  Ser Leu Ala Ala Cys  Leu Val Val Glu Ala  Val Val Trp
437     1025            1030            1035
440 Lys Ser  Val Thr Lys Asn Arg  Thr Ser Tyr Met Arg  His Thr Cys
441     1040            1045            1050
444 Ile Val  Asn Ile Ala Ala Ser  Leu Leu Val Ala Asn  Thr Trp Phe
445     1055            1060            1065
448 Ile Val  Val Ala Ala Ile Gln  Asp Asn Arg Tyr Ile  Leu Cys Lys
449     1070            1075            1080
452 Thr Ala  Cys Val Ala Ala Thr  Phe Phe Ile His Phe  Phe Tyr Leu
453     1085            1090            1095
456 Ser Val  Phe Phe Trp Met Leu  Thr Leu Gly Leu Met  Leu Phe Tyr
457     1100            1105            1110
460 Arg Leu  Val Phe Ile Leu His  Glu Thr Ser Arg Ser  Thr Gln Lys
461     1115            1120            1125
464 Ala Ile  Ala Phe Cys Leu Gly  Tyr Gly Cys Pro Leu  Ala Ile Ser
465     1130            1135            1140
468 Val Ile  Thr Leu Gly Ala Thr  Gln Pro Arg Glu Val  Tyr Thr Arg
469     1145            1150            1155
472 Lys Asn  Val Cys Trp Leu Asn  Trp Glu Asp Thr Lys  Ala Leu Leu
473     1160            1165            1170
476 Ala Phe  Ala Ile Pro Ala Leu  Ile Ile Val Val Val  Asn Ile Thr
477     1175            1180            1185
480 Ile Thr  Ile Val Val Ile Thr  Lys Ile Leu Arg Pro  Ser Ile Gly
481     1190            1195            1200
484 Asp Lys  Pro Cys Lys Gln Glu  Lys Ser Ser Leu Phe  Gln Ile Ser
485     1205            1210            1215
488 Lys Ser  Ile Gly Val Leu Thr  Pro Leu Leu Gly Leu  Thr Trp Gly
489     1220            1225            1230
492 Phe Gly  Leu Thr Thr Val Phe  Pro Gly Thr Asn Leu  Val Phe His
493     1235            1240            1245
496 Ile Ile  Phe Ala Ile Leu Asn  Val Phe Gln Gly Leu  Phe Ile Leu
497     1250            1255            1260
500 Leu Phe  Gly Cys Leu Trp Asp  Leu Lys Val Gln Glu  Ala Leu Leu
```

RAW SEQUENCE LISTING   DATE: 12/30/2005
PATENT APPLICATION: US/10/561,132   TIME: 09:22:52

Input Set : A:\60.US2.PCT.txt
Output Set: N:\CRF4\12302005\J561132.raw

```
501     1265               1270               1275
504 Asn Lys  Phe Ser Leu Ser Arg  Trp Ser Ser Gln His  Ser Lys Ser
505     1280               1285               1290
508 Thr Ser  Leu Gly Ser Ser Thr  Pro Val Phe Ser Met  Ser Ser Pro
509     1295               1300               1305
512 Ile Ser  Arg Arg Phe Asn Asn  Leu Phe Gly Lys Thr  Gly Thr Tyr
513     1310               1315               1320
516 Asn Val  Ser Thr Pro Glu Ala  Thr Ser Ser Ser Leu  Glu Asn Ser
517     1325               1330               1335
520 Ser Ser  Ala Ser Ser Leu Leu  Asn
521     1340               1345
```

RAW SEQUENCE LISTING ERROR SUMMARY   DATE: 12/30/2005
PATENT APPLICATION: US/10/561,132   TIME: 09:22:53

Input Set : **A:\60.US2.PCT.txt**
Output Set: **N:\CRF4\12302005\J561132.raw**

Invalid <213> Response:

**Use of "Artificial" only as "<213> Organism" response is incomplete,
per 1.823(b) of New Sequence Rules. Valid response is Artificial Sequence.**

Seq#:7,8,9,10,11,12

VERIFICATION SUMMARY   DATE: 12/30/2005
PATENT APPLICATION: US/10/561,132   TIME: 09:22:53

Input Set : **A:\60.US2.PCT.txt**
Output Set: **N:\CRF4\12302005\J561132.raw**

```
L:11 M:270 C: Current Application Number differs, Replaced Current Application No
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:170 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:174 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
L:178 M:200 E: Mandatory Header Field missing, <220> Tag not found for SEQ ID#:2
L:178 M:258 W: Mandatory Feature missing, <220> Tag not found for SEQ ID#:2
```